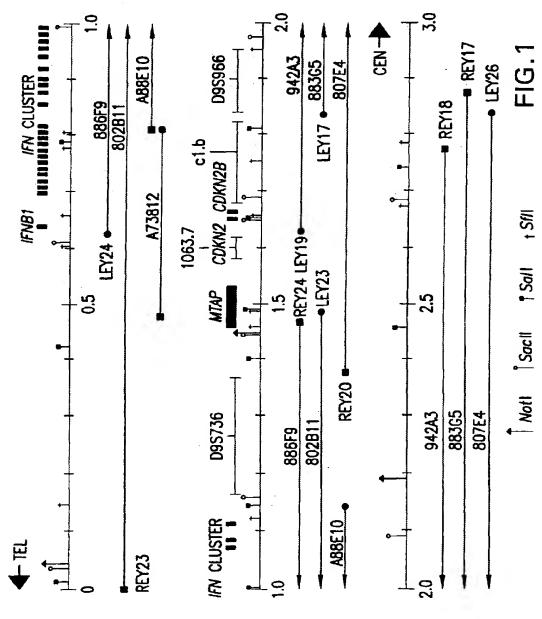


7IG. 19E







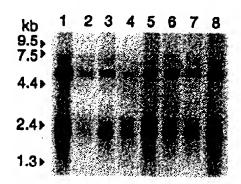


FIG. 2A



FIG. 2B



```
1 gaatteeget cegeactget cacteeegeg cagtgaggtt ggcacageea eegetetgtg
  61 getegettgg ttecettagt ecegageget egeceaetge agatteettt ecegtgeaga
 121 categoratet ggcaccacca coacegorgt gaagattgga ataattggtg gaacaggoot
 181 ggatgatcca gaaattttag aaggaagaac tgaaaaatat gtggatactc catttggcaa
 241 gccatctgat gccttaattt tggggaagat aaaaaatgtt gattgcatcc tccttgcaag
 301 gcatggaagg cagcacacca tcatgccttc aaaggtcaaa taccaggcga acatctgggc
 361 tttgaaggaa gagggctgta cacatgtcat agtgaccaca gattgtggct ccttgaggga
 421 ggagattcag cccggcgata ttgtcattat tgatcagttc attgacagga ccactatgag
 481 acctcagtcc ttctatgatg gaagtcattc ttgtgccaga ggagtgtgcc atattccaat
 541 ggatgagccg ttttgcccca aaacgagaga ggttcttata gagactgcta agaagctagg
 601 actccggtgc cactcaaagg ggacaatggt cacaatcgag ggacctcgtt ttagctccag
 661 ggcagaaagc ttcatgttcc gcacctgggg ggcggatgtt atcaacatga ccacagttcc
 721 agaggtggtt cttgctaagg aggctggaat ttgttacgca agtatcgcca tggcgacaga
 781 ttatgactgc tggaaggagc acgaggaagc agtttcggtg gaccgggtct taaagaccct
 841 gaaagaaaac gctaataaag ccaaaagctt actgctcact accatacctc agatagggtc
 901 cacagaatgg tcagaaaccc tccataacct gaagaatatg gcccagtttt ctgttttatt
 961 accaagacat taaagtagca tggctgccca ggagaaaaga agacattcta attccagtca
1021 ttttgggaat tcctgcttaa cttgaaaaaa atatgggaaa gacatgcagc tttcatgccc
1081 ttgcctatca aagagtatgt tgtaagaaag acaagacatt gtgtgtatta gagactcctg
1141 aatgatttag acaacttcaa aatacagaag aaaagcaaat gactagtaaa catgtgggaa
1261 atttqcaaca ataaagggtg gagggtaatc tctactttcc tatactgcca aagaatgtga
1321 ggaagaaatg ggactctttg gttatttatt gatgcgactg taaattggta cagtatttct
1381 ggagggcaat ttggtaaaat gcatcaaaag acttaaaaat acggacgtcc tttggtgctg
1441 qqaaatctac atataqcaat ttctctttaa aaccatatca gagatgcata caaagaatta
1501 tatataaaga agggtgttta ataatgatag ttataataat aaataattga aacaatctga
1561 atcccttgca attggaggta aattatgtct tagttataat ctagattgtg aatcagccaa
1621 ctgaaaatcc tttttgcata tttcaatgtc ctaaaaagac acggttgctc tatatatgaa
1681 gtgaaaaaag gatatggtag cattttatag tactagtttt gctttaaaat gctatgtaaa
1741 tatacaaaaa aactagaaag aaatatatat aaccttgtta ttgtatttgg gggagggata
1801 ctgqqataat ttttattttc tttgaatctt tctgtgtctt cacatttttc tacagtgaat
1861 ataatcaaat agtaaagggc cgtaaaaata aaagtggatt tagaaagatc cagttcttga
1921 aaacactgtt tctggtaatg aagcagaatt taagttggta atattaaggt gaatgtcatt
1981 taagggagtt acatetttat tetgetaaag aagaggatea tigatitetg tacagteaga
2041 acagtacttg ggtgtgcaac agctttctga gaaaagctag gtgtataata gtttaactga
2101 aagtttaact atttaaaaga ctaaatgcac attttatggt atctgatatt ttaaaaagta
2161 atgtgagett eteettitta tgagttaaat tattitatae gagttggtaa titgtgeett
2221 ttaataaagt ggaagettge tttttaaaaa aaaaaaaaa geggaatte
```



1	MASGTTTTAVKIGIIGGTGLDDPEILEGRTEKYVDTPFGK	40
41	PSDALILGKIKNVDCILLARHGRQHTIMPSKVNYQANIWA	80
81	LKEEGCTHVIVTTACGSLREEIQPGDIVIIDQFIDRTTMR	120
121	PQSFYDGSHSCARGVCHIPMAEPFCPKTREVLIETAKKLG	160
161	LRCHSKGTMVTIEGPRFSSRAESFMFRTWGADVINMTTVP	200
201	EVVLAKEAGICYASIAMATDYDCWKEHEEAVSVDRVLKTL	240
241	KENANKAKSLLLTTIPQIGSTEWSETLHNLKNMAQFSVLL	280
281	PRH 283	

FIG.3B



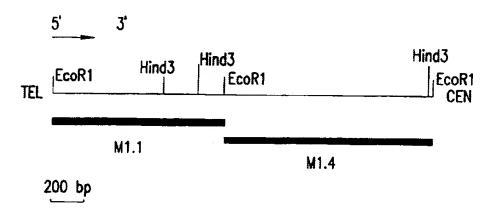


FIG.4



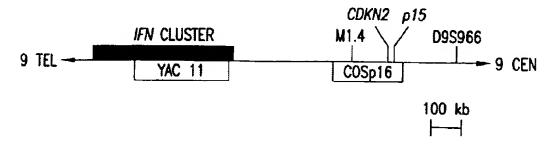


FIG.5





FIG. 6A



FIG. 6B



FIG. 6C



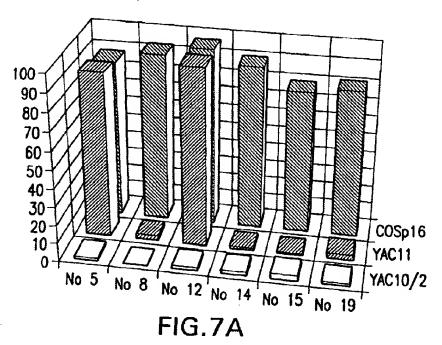
FIG. 6D

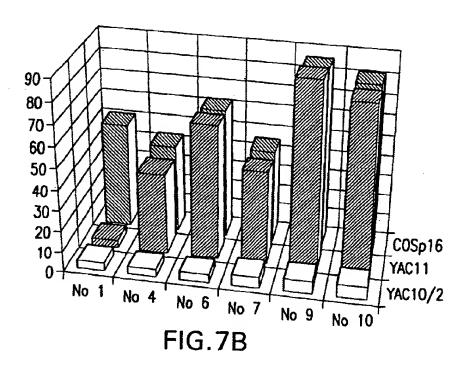




FIG. 6E









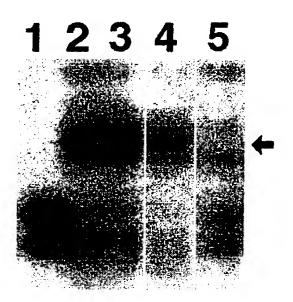
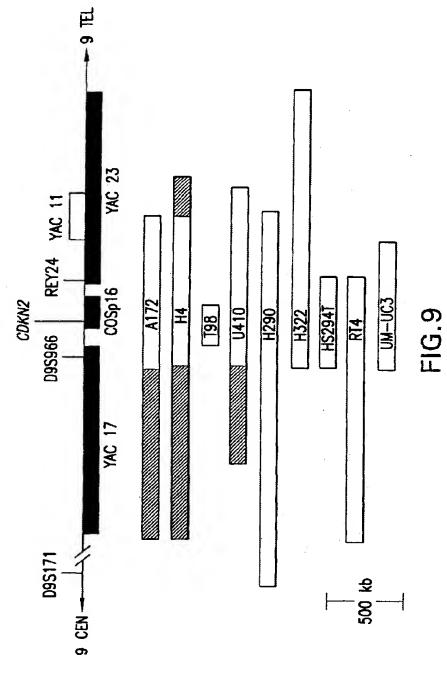
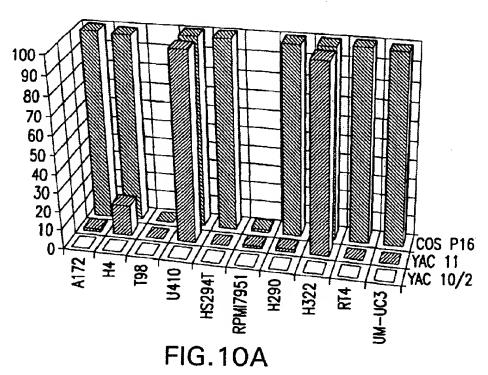


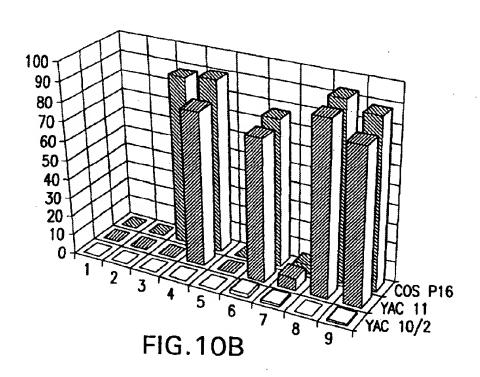
FIG. 8













METHYLTHIOADENOSINE PHOSPHORYLASE CDNA MAP

CENTROMERE	Hind Hind								
1ELUMERE 5' 3'	INITIATION SITE	EXONS: 1 2 3 4 5 6 7	PRIMERS: AF AR BF BR OF DF EF FF GF, FR GR HF HR	PROBE PROBE PROBE DROBE A BF-BR C C E	PROBE PROBE PROBE GF-GR AF-AR CF-CR FF-FR FF-FR	pM1.1	cosmid 65G7:	too bp cosmid 22588:	FIG



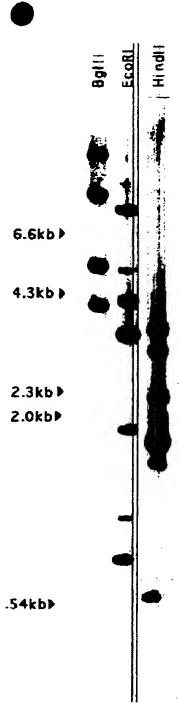


FIG. 12





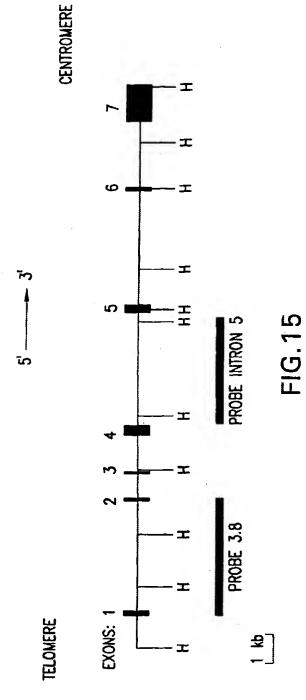
FIG. 13

34kd) 🗰

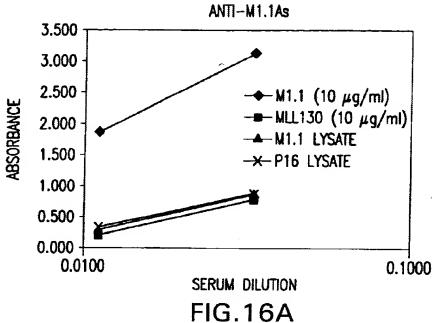
FIG. 14

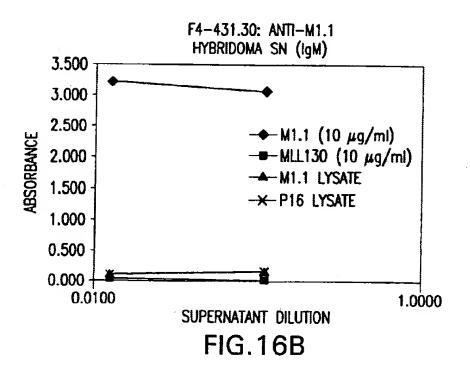


GENOMIC ORGANIZATION OF THE METHYLTHIOADENOSINE PHOSPHORYLASE GENE

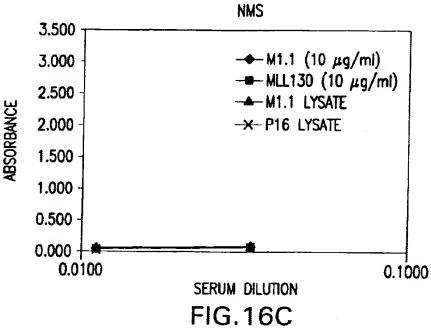


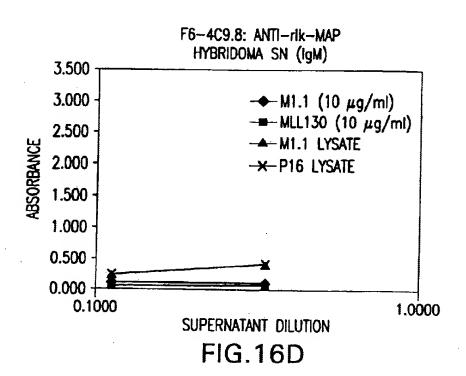














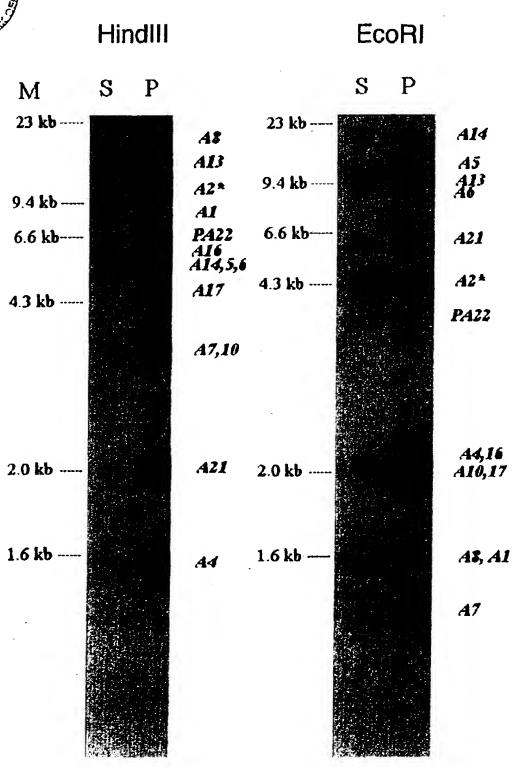


FIG. 17A



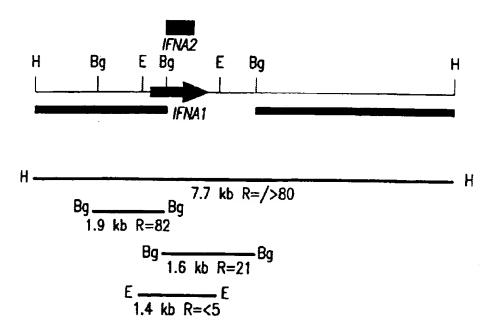
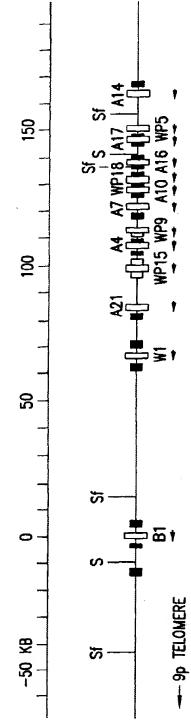
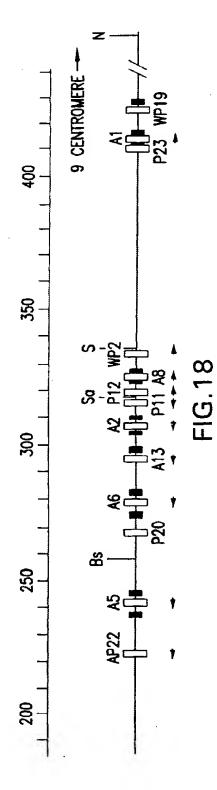
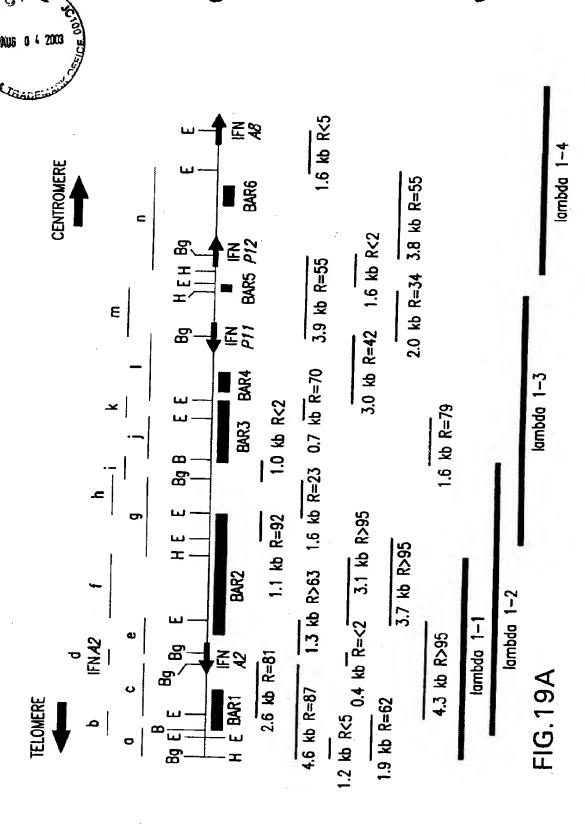


FIG.17B











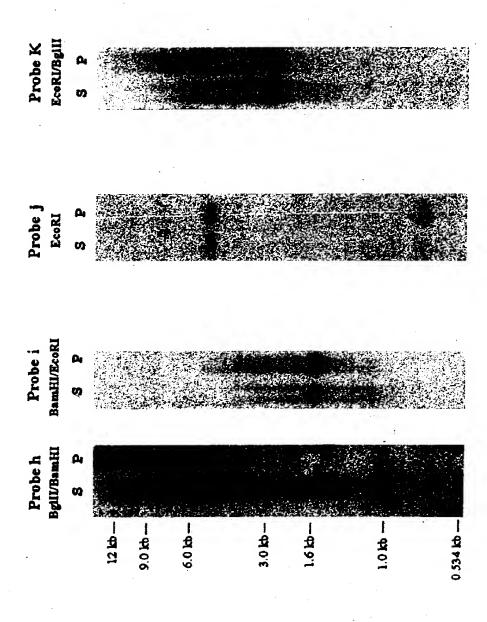
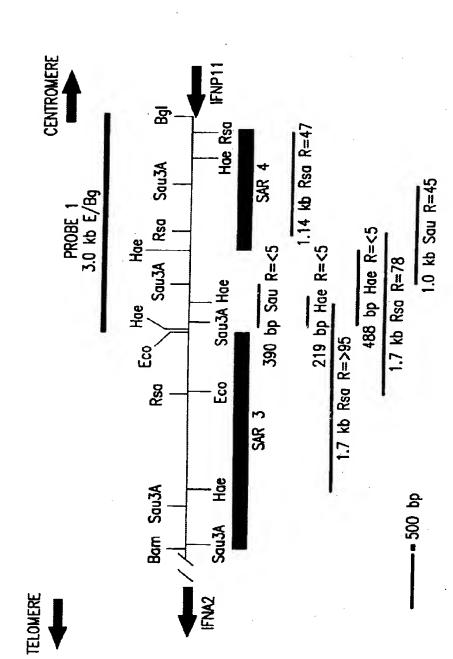


FIG. 19B



1.0 kb Hae R=42

FIG.20



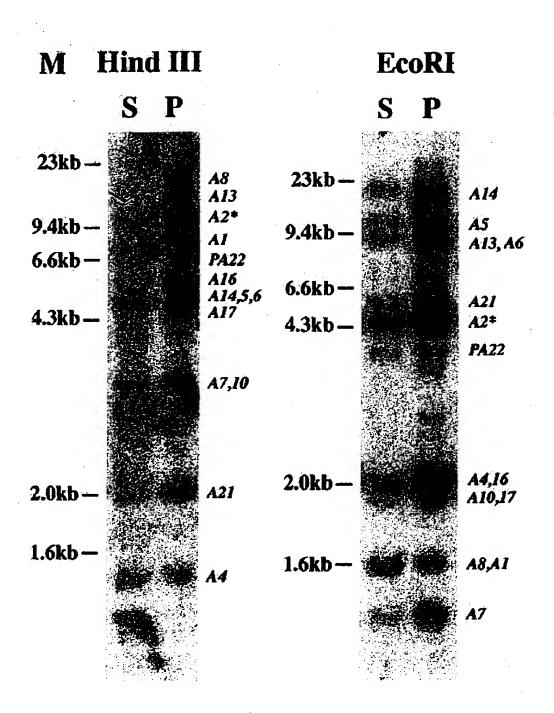


FIG. 21